IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANT: MING-MING ZHOU *ET AL.*

SERIAL NO. :   UNASSIGNED   EXAMINER :   UNKNOWN

FILED   :   HEREWITH   ART UNIT   :   UNKNOWN

FOR   :   METHODS OF IDENTIFYING MODULATORS OF
                   BROMODOMAINS

EXPRESS MAIL "MAILING LABEL NO."   :   EL676518587US
DATE OF DEPOSIT   :   February 16, 2001

STATEMENT IN SUPPORT OF THE FILING/SUBMISSION OF A NUCLEOTIDE/AMINO ACID SEQUENCE LISTING IN ACCORDANCE WITH 37 CFR §1.821 - 1.825

ASSISTANT COMMISSIONER FOR PATENTS
WASHINGTON, D.C. 20231

Dear Sir:

MICHAEL D. DAVIS, attorney of record, hereby states as follows:

I hereby state that the content of the paper and computer readable copies of the Sequence Listing submitted in accordance with 37 CFR §1.821(c), (e) and (f) respectively, are the same.

DATED: February 16, 2001

_____
MICHAEL D. DAVIS

SEQUENCE LISTING

<110> Zhou, Ming-Ming
      Aggarwal, Aneel
      Verdin, Eric
      Ott, Melanie

<120> Methods of Identifying Modulators of Bromodomains

<130> 2459-1-003CIP

<140> Unassigned
<141> 2001-02-16

<150> 09/510,314
<151> 2000-02-22

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<170> PatentIn version 3.0

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<212> PRT
<213> Homo sapiens

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            35   40   45

Ser Gly Ala Cys Gly Pro Ala Thr Ala Val Ala Ala Ala Gly Thr Ala
        50   55   60

Glu Gly Pro Gly Gly Gly Gly Ser Ala Arg Ile Ala Val Lys Lys Ala
65   70   75   80

Gln Leu Arg Ser Ala Pro Arg Ala Lys Lys Leu Glu Lys Leu Gly Val
                85   90   95

Tyr Ser Ala Cys Lys Ala Glu Glu Ser Cys Lys Cys Asn Gly Trp Lys

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                100                   105                   110
Asn Pro Asn Pro Ser Pro Thr Pro Pro Arg Ala Asp Leu Gln Gln Ile
        115             120             125
Ile Val Ser Leu Thr Glu Ser Cys Arg Ser Cys Ser His Ala Leu Ala
    130             135             140
Ala His Val Ser His Leu Glu Asn Val Ser Glu Glu Glu Met Asn Arg
145             150             155             160
Leu Leu Gly Ile Val Leu Asp Val Glu Tyr Leu Phe Thr Cys Val His
            165             170             175
Lys Glu Glu Asp Ala Asp Thr Lys Gln Val Tyr Phe Tyr Leu Phe Lys
        180             185             190
Leu Leu Arg Lys Ser Ile Leu Gln Arg Gly Lys Pro Val Val Glu Gly
        195             200             205
Ser Leu Glu Lys Lys Pro Pro Phe Glu Lys Pro Ser Ile Glu Gln Gly
    210             215             220
Val Asn Asn Phe Val Gln Tyr Lys Phe Ser His Leu Pro Ala Lys Glu
225             230             235             240
Arg Gln Thr Ile Val Glu Leu Ala Lys Met Phe Leu Asn Arg Ile Asn
            245             250             255
Tyr Trp His Leu Glu Ala Pro Ser Gln Arg Arg Leu Arg Ser Pro Asn
            260             265             270
Asp Asp Ile Ser Gly Tyr Lys Glu Asn Tyr Thr Arg Trp Leu Cys Tyr
        275             280             285
Cys Asn Val Pro Gln Phe Cys Asp Ser Leu Pro Arg Tyr Glu Thr Thr
        290             295             300
Gln Val Phe Gly Arg Thr Leu Leu Arg Ser Val Phe Thr Val Met Arg
305             310             315             320
Arg Gln Leu Leu Glu Gln Ala Arg Gln Glu Lys Asp Lys Leu Pro Leu
            325             330             335
Glu Lys Arg Thr Leu Ile Leu Thr His Phe Pro Lys Phe Leu Ser Met
            340             345             350
Leu Glu Glu Glu Val Tyr Ser Gln Asn Ser Pro Ile Trp Asp Gln Asp
        355             360             365
Phe Leu Ser Ala Ser Ser Arg Thr Ser Gln Leu Gly Ile Gln Thr Val
        370             375             380
Ile Asn Pro Pro Pro Val Ala Gly Thr Ile Ser Tyr Asn Ser Thr Ser
385             390             395             400
Ser Ser Leu Glu Gln Pro Asn Ala Gly Ser Ser Ser Pro Ala Cys Lys
            405             410             415
Ala Ser Ser Gly Leu Glu Ala Asn Pro Gly Glu Lys Arg Lys Met Thr
            420             425             430
Asp Ser His Val Leu Glu Glu Ala Lys Lys Pro Arg Val Met Gly Asp
        435             440             445
Ile Pro Met Glu Leu Ile Asn Glu Val Met Ser Thr Ile Thr Asp Pro
    450             455             460
Ala Ala Met Leu Gly Pro Glu Thr Asn Phe Leu Ser Ala His Ser Ala
```

```
        465                    470                    475                    480
        Arg Asp Glu Ala Ala Arg Leu Glu Glu Arg Arg Gly Val Ile Glu Phe
                        485                    490                    495

        His Val Val Gly Asn Ser Leu Asn Gln Lys Pro Asn Lys Lys Ile Leu
                        500                    505                    510

        Met Trp Leu Val Gly Leu Gln Asn Val Phe Ser His Gln Leu Pro Arg
                        515                    520                    525

        Met Pro Lys Glu Tyr Ile Thr Arg Leu Val Phe Asp Pro Lys His Lys
                530                    535                    540

        Thr Leu Ala Leu Ile Lys Asp Gly Arg Val Ile Gly Gly Ile Cys Phe
        545                    550                    555                    560

        Arg Met Phe Pro Ser Gln Gly Phe Thr Glu Ile Val Phe Cys Ala Val
                        565                    570                    575

        Thr Ser Asn Glu Gln Val Lys Gly Tyr Gly Thr His Leu Met Asn His
                        580                    585                    590

        Leu Lys Glu Tyr His Ile Lys His Asp Ile Leu Asn Phe Leu Thr Tyr
                        595                    600                    605

        Ala Asp Glu Tyr Ala Ile Gly Tyr Phe Lys Lys Gln Gly Phe Ser Lys
                610                    615                    620

        Glu Ile Lys Ile Pro Lys Thr Lys Tyr Val Gly Tyr Ile Lys Asp Tyr
        625                    630                    635                    640

        Glu Gly Ala Thr Leu Met Gly Cys Glu Leu Asn Pro Arg Ile Pro Tyr
                        645                    650                    655

        Thr Glu Phe Ser Val Ile Ile Lys Lys Gln Lys Glu Ile Ile Lys Lys
                        660                    665                    670

        Leu Ile Glu Arg Lys Gln Ala Gln Ile Arg Lys Val Tyr Pro Gly Leu
                        675                    680                    685

        Ser Cys Phe Lys Asp Gly Val Arg Gln Ile Pro Ile Glu Ser Ile Pro
                        690                    695                    700

        Gly Ile Arg Glu Thr Gly Trp Lys Pro Ser Gly Lys Glu Lys Ser Lys
        705                    710                    715                    720

        Glu Pro Arg Asp Pro Asp Gln Leu Tyr Ser Thr Leu Lys Ser Ile Leu
                        725                    730                    735

        Gln Gln Val Lys Ser His Gln Ser Ala Trp Pro Phe Met Glu Pro Val
                        740                    745                    750

        Lys Arg Thr Glu Ala Pro Gly Tyr Tyr Glu Val Ile Arg Phe Pro Met
                        755                    760                    765

        Asp Leu Lys Thr Met Ser Glu Arg Leu Lys Asn Arg Tyr Tyr Val Ser
                770                    775                    780

        Lys Lys Leu Phe Met Ala Asp Leu Gln Arg Val Phe Thr Asn Cys Lys
        785                    790                    795                    800

        Glu Tyr Asn Ala Ala Glu Ser Glu Tyr Tyr Lys Cys Ala Asn Ile Leu
                        805                    810                    815

        Glu Lys Phe Phe Phe Ser Lys Ile Lys Glu Ala Gly Leu Ile Asp Lys
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<211>   12
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<213>   artificial sequence

<220>
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<222>   (2)..(2)
<223>   X is two to three amino acids.  Each of these can be any amino
ac
        id


<220>
<221>   X
<222>   (4)..(4)
<223>   The X is five to eight amino acids.  Each of these can be any
ami
        no acid


<220>
<221>   X
<222>   (5)..(5)
<223>   X is a single amino acid that is either Pro, Lys, or His.


<220>
<221>   X
<222>   (6)..(6)
<223>   This X is any single amino acid.


<220>
<221>   X
<222>   (8)..(8)
<223>   This X is a single amino acid that can be either Tyr, Phe, or
His


<220>
<221>   X
<222>   (9)..(9)
<223>   X is 5 amino  acids.  Each of these can be any amino acid.


<220>
<221>   X
<222>   (11)..(11)
<223>   X is a single amino acid that can be Met, Ile, or Val.


<400>   3

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<210>   4
<211>   12
<212>   PRT
<213>   artificial sequence

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<222>   (6)..(6)
<223>   The X represents an acetyl-lysine


<400>   4
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Ile Ser Tyr Gly Arg Xaa Lys Arg Arg Gln Arg Arg
1               5                   10
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<210>   5
<211>   14
<212>   PRT
<213>   artificial sequence
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<220>
<221>   X
<222>   (8)..(8)
<223>   The X represents an acetyl lysine.
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```
<400>   5
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Ala Arg Lys Ser Thr Gly Gly Xaa Ala Pro Arg Lys Gln Leu
1               5                   10
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<210>   6
<211>   14
<212>   PRT
<213>   artificial sequence
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<220>
<221>   X
<222>   (8)..(8)
<223>   The X represents an acetyl lysine.
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<400>   6
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1               5                   10
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<210>   7
<211>   110
<212>   PRT
<213>   Homo sapiens, peptide
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```
<400>   7
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Ser Lys Glu Pro Arg Asp Pro Asp Gln Leu Tyr Ser Thr Leu Lys Ser
1               5                   10                  15

Ile Leu Gln Gln Val Lys Ser His Gln Ser Ala Trp Pro Phe Met Glu
            20                  25                  30

Pro Val Lys Arg Thr Glu Ala Pro Gly Tyr Tyr Glu Val Ile Arg Ser
            35                  40                  45

Pro Met Asp Leu Lys Thr Met Ser Glu Arg Leu Lys Asn Arg Tyr Tyr
        50                  55                  60

Val Ser Lys Lys Leu Phe Met Ala Asp Leu Gln Arg Val Phe Thr Asn
65                  70                  75                  80

Cys Lys Glu Tyr Asn Ala Pro Glu Ser Glu Tyr Tyr Lys Cys Ala Asn
                85                  90                  95

Ile Leu Glu Lys Phe Phe Phe Ser Lys Ile Lys Glu Ala Gly
            100                 105                 110
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<210>   8
<211>   110
<212>   PRT
<213>   Homo sapiens
```

<400> 8

Gly Lys Glu Leu Lys Asp Pro Asp Gln Leu Tyr Thr Thr Leu Lys Asn
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Leu Leu Ala Gln Ile Lys Ser His Pro Ser Ala Trp Pro Phe Met Glu
                20   25   30

Pro Val Lys Lys Ser Glu Ala Pro Asp Tyr Tyr Glu Val Ile Arg Phe
            35   40   45

Pro Ile Asp Leu Lys Thr Met Thr Glu Arg Leu Arg Ser Arg Tyr Tyr
        50   55   60

Val Thr Arg Lys Leu Phe Val Ala Asp Leu Gln Arg Val Ile Ala Asn
65   70   75   80

Cys Arg Glu Tyr Asn Pro Pro Asp Ser Glu Tyr Cys Arg Cys Ala Ser
                85   90   95

Ala Leu Glu Lys Phe Phe Tyr Phe Lys Leu Lys Glu Gly Gly
            100   105   110

<210>   9
<211>   109
<212>   PRT
<213>   Tetrahymena thermophila

<400>   9

Leu Lys Lys Ser Lys Glu Arg Ser Phe Asn Leu Gln Cys Ala Asn Val
1   5   10   15

Ile Glu Asn Met Lys Arg His Lys Gln Ser Trp Pro Phe Leu Asp Pro
                20   25   30

Val Asn Lys Asp Asp Val Pro Asp Tyr Tyr Asp Val Ile Thr Asp Pro
            35   40   45

Ile Asp Ile Lys Ala Ile Glu Lys Lys Leu Gln Asn Asn Gln Tyr Val
        50   55   60

Asp Lys Asp Gln Phe Ile Lys Asp Val Lys Arg Ile Phe Thr Asn Ala
65   70   75   80

Lys Ile Tyr Asn Gln Pro Asp Thr Ile Tyr Tyr Lys Ala Ala Lys Glu
                85   90   95

Leu Glu Asp Phe Val Glu Pro Tyr Leu Thr Lys Leu Lys
            100   105

<210>   10
<211>   109
<212>   PRT
<213>   Saccharomyces cerevisiae

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Ala Gln Arg Pro Lys Arg Gly Pro His Asp Ala Ala Ile Gln Asn Ile
1   5   10   15

Leu Thr Glu Leu Gln Asn His Ala Ala Ala Trp Pro Phe Leu Gln Pro
                20   25   30

Val Asn Lys Glu Glu Val Pro Asp Tyr Tyr Asp Phe Ile Lys Glu Pro
            35   40   45

Met Asp Leu Ser Thr Met Glu Ile Lys Leu Glu Ser Asn Lys Tyr Gln
        50   55   60

```
Lys Met Glu Asp Phe Ile Tyr Asp Ala Arg Leu Val Phe Asn Asn Cys
65              70              75              80

Arg Met Tyr Asn Gly Glu Asn Thr Ser Tyr Tyr Lys Tyr Ala Asn Arg
            85              90              95

Leu Glu Lys Phe Phe Asn Asn Lys Val Lys Glu Ile Pro
            100             105

<210>   11
<211>   112
<212>   PRT
<213>   Homo sapiens

<400>   11

Lys Lys Ile Phe Lys Pro Glu Glu Leu Arg Gln Ala Leu Met Pro Thr
1               5               10              15

Leu Glu Ala Leu Tyr Arg Gln Asp Pro Glu Ser Leu Pro Phe Arg Gln
            20              25              30

Pro Val Asp Pro Gln Leu Leu Gly Ile Pro Asp Tyr Phe Asp Ile Val
            35              40              45

Lys Ser Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly
            50              55              60

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Ile Trp Leu Met Phe
65              70              75              80

Asn Asn Ala Trp Leu Tyr Asn Arg Lys Thr Ser Arg Val Tyr Lys Tyr
            85              90              95

Cys Ser Lys Leu Ser Glu Val Phe Glu Gln Glu Ile Asp Pro Val Met
            100             105             110

<210>   12
<211>   112
<212>   PRT
<213>   Homo sapiens

<400>   12

Lys Lys Ile Phe Lys Pro Glu Glu Leu Arg Gln Ala Leu Met Pro Thr
1               5               10              15

Leu Glu Ala Leu Tyr Arg Gln Asp Pro Glu Ser Leu Pro Phe Arg Gln
            20              25              30

Pro Val Asp Pro Gln Leu Leu Gly Ile Pro Asp Tyr Phe Asp Ile Val
            35              40              45

Lys Asn Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly
            50              55              60

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Val Trp Leu Met Phe
65              70              75              80

Asn Asn Ala Trp Leu Tyr Asn Arg Lys Thr Ser Arg Val Tyr Lys Phe
            85              90              95

Cys Ser Lys Leu Ala Glu Val Phe Glu Gln Glu Ile Asp Pro Val Met
            100             105             110

<210>   13
<211>   112
<212>   PRT
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<213> Mus musculus

<400> 13

Lys Lys Ile Phe Lys Pro Glu Glu Leu Arg Gln Ala Leu Met Pro Thr
1   5   10   15

Leu Glu Ala Leu Tyr Arg Gln Asp Pro Glu Ser Leu Pro Phe Arg Gln
            20   25   30

Pro Val Asp Pro Gln Leu Leu Gly Ile Pro Asp Tyr Phe Asp Ile Val
        35   40   45

Lys Asn Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly
    50   55   60

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Val Arg Leu Met Phe
65   70   75   80

Asn Asn Ala Trp Leu Tyr Asn Arg Lys Thr Ser Arg Val Tyr Lys Phe
                85   90   95

Cys Ser Lys Leu Ala Glu Val Phe Glu Gln Glu Ile Asp Pro Val Met
                100   105   110

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<211> 111
<212> PRT
<213> Caenorhabditis elegans

<400> 14

Asp Thr Val Phe Ser Gln Glu Asp Leu Ile Lys Phe Leu Leu Pro Val
1   5   10   15

Trp Glu Lys Leu Asp Lys Ser Glu Asp Ala Ala Pro Phe Arg Val Pro
            20   25   30

Val Asp Ala Lys Leu Leu Asn Ile Pro Asp Tyr His Glu Ile Ile Lys
        35   40   45

Arg Pro Met Asp Leu Glu Thr Val His Lys Lys Leu Tyr Ala Gly Gln
    50   55   60

Tyr Gln Asn Ala Gly Gln Phe Cys Asp Asp Ile Trp Leu Met Leu Asp
65   70   75   80

Asn Ala Trp Leu Tyr Asn Arg Lys Asn Ser Lys Val Tyr Lys Tyr Gly
                85   90   95

Leu Lys Leu Ser Glu Met Phe Val Ser Glu Met Asp Pro Val Met
                100   105   110

<210> 15
<211> 110
<212> PRT
<213> Homo sapiens

<400> 15

Arg Arg Arg Thr Asp Pro Met Val Thr Leu Ser Ser Ile Leu Glu Ser
1   5   10   15

Ile Ile Asn Asp Met Arg Asp Leu Pro Asn Thr Tyr Pro Phe His Thr
            20   25   30

Pro Val Asn Ala Lys Val Val Lys Asp Tyr Tyr Lys Ile Ile Thr Arg
        35   40   45

```
Pro Met Asp Leu Gln Thr Leu Arg Glu Asn Val Arg Lys Arg Leu Tyr
    50              55              60
Pro Ser Arg Glu Glu Phe Arg Glu His Leu Glu Leu Ile Val Lys Asn
65              70              75              80
Ser Ala Thr Tyr Asn Gly Pro Lys His Ser Leu Thr Gln Ile Ser Gln
                85              90              95
Ser Met Leu Asp Leu Cys Asp Glu Lys Leu Lys Glu Lys Glu
            100             105             110
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<210> 16
<211> 110
<212> PRT
<213> Mesocricetus auratus

<400> 16

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1               5               10              15
Ile Ile Asn Asp Met Arg Asp Leu Pro Asn Thr Tyr Pro Phe His Thr
            20              25              30
Pro Val Asn Ala Lys Val Val Lys Asp Tyr Tyr Lys Ile Ile Thr Arg
            35              40              45
Pro Met Asp Leu Gln Thr Leu Arg Glu Asn Val Arg Lys Arg Leu Tyr
    50              55              60
Pro Ser Arg Glu Glu Phe Arg Glu His Leu Glu Leu Ile Val Lys Asn
65              70              75              80
Ser Ala Thr Tyr Asn Gly Pro Lys His Ser Leu Thr Gln Ile Ser Gln
                85              90              95
Ser Met Leu Asp Leu Cys Asp Glu Lys Leu Lys Glu Lys Glu
            100             105             110
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<210> 17
<211> 111
<212> PRT
<213> Homo sapiens

<400> 17

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Ile Val Thr Gln Lys Met Met Ala Val Pro Asp Ser Trp Pro Phe His
            20              25              30
His Pro Val Asn Lys Lys Phe Val Pro Asp Tyr Tyr Lys Val Ile Val
        35              40              45
Asn Pro Met Asp Leu Glu Thr Ile Arg Lys Asn Ile Ser Lys His Lys
    50              55              60
Tyr Gln Ser Arg Glu Ser Phe Leu Asp Asp Val Asn Leu Ile Leu Ala
65              70              75              80
Asn Ser Val Lys Tyr Asn Gly Pro Glu Ser Gln Tyr Thr Lys Thr Ala
                85              90              95
Gln Glu Ile Val Asn Val Cys Tyr Gln Thr Leu Thr Glu Tyr Asp
            100             105             110
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<210> 18
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<211>    111
<212>    PRT
<213>    Mesocricetus auratus

<400>    18

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1               5                   10                  15

Ile Val Thr Gln Lys Met Met Ala Val Pro Asp Ser Trp Pro Phe His
            20                  25                  30

His Pro Val Asn Lys Lys Phe Val Pro Asp Tyr Tyr Lys Val Ile Val
        35                  40                  45

Ser Pro Met Asp Leu Glu Thr Ile Arg Lys Asn Ile Ser Lys His Lys
    50                  55                  60

Tyr Gln Ser Arg Glu Ser Phe Leu Asp Asp Val Asn Leu Ile Leu Ala
65                  70                  75                  80

Asn Ser Val Lys Tyr Asn Gly Ser Glu Ser Gln Tyr Thr Lys Thr Ala
                85                  90                  95

Gln Glu Ile Val Asn Val Cys Tyr Gln Thr Leu Thr Glu Tyr Asp
            100                 105                 110

<210>    19
<211>    111
<212>    PRT
<213>    Homo sapiens

<400>    19

Lys Pro Gly Arg Val Thr Asn Gln Leu Gln Tyr Leu His Lys Val Val
1               5                   10                  15

Met Lys Ala Leu Trp Lys His Gln Phe Ala Trp Pro Phe Arg Gln Pro
            20                  25                  30

Val Asp Ala Val Lys Leu Gly Leu Pro Asp Tyr His Lys Ile Ile Lys
        35                  40                  45

Gln Pro Met Asp Met Gly Thr Ile Lys Arg Arg Leu Glu Asn Asn Tyr
    50                  55                  60

Tyr Trp Ala Ala Ser Glu Cys Met Gln Asp Phe Asn Thr Met Phe Thr
65                  70                  75                  80

Asn Cys Tyr Ile Tyr Asn Lys Pro Thr Asp Asp Ile Val Leu Met Ala
                85                  90                  95

Gln Thr Leu Glu Lys Ile Phe Leu Gln Lys Val Ala Ser Met Pro
            100                 105                 110

<210>    20
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<213>    Homo sapiens

<400>    20

Lys Pro Gly Arg Lys Thr Asn Gln Leu Gln Tyr Met Gln Asn Val Val
1               5                   10                  15

Val Lys Thr Leu Trp Lys His Gln Phe Ala Trp Pro Phe Tyr Gln Pro
            20                  25                  30

Val Asp Ala Ile Lys Leu Asn Leu Pro Asp Tyr His Lys Ile Ile Lys
```

```
            35                    40                    45
Asn Pro Met Asp Met Gly Thr Ile Lys Lys Arg Leu Glu Asn Asn Tyr
    50                  55                  60

Tyr Trp Ser Ala Ser Glu Cys Met Gln Asp Phe Asn Thr Met Phe Thr
65                  70                  75                  80

Asn Cys Tyr Ile Tyr Asn Lys Pro Thr Asp Asp Ile Val Leu Met Ala
                85                  90                  95

Gln Ala Leu Glu Lys Ile Phe Leu Gln Lys Val Ala Gln Met Pro
            100                 105                 110
```

```
<210>   21
<211>   111
<212>   PRT
<213>   Drosophila melanogaster

<400>   21
```

```
Arg Pro Gly Arg Asn Thr Asn Gln Leu Gln Tyr Leu Ile Lys Thr Val
1               5                   10                  15

Met Lys Val Ile Trp Lys His His Phe Ser Trp Pro Phe Gln Gln Pro
                20                  25                  30

Val Asp Ala Lys Lys Leu Asn Leu Pro Asp Tyr His Lys Ile Ile Lys
        35                  40                  45

Gln Pro Met Asp Met Gly Thr Ile Lys Lys Arg Leu Glu Asn Asn Tyr
    50                  55                  60

Tyr Trp Ser Ala Lys Glu Thr Ile Gln Asp Phe Asn Thr Met Phe Asn
65                  70                  75                  80

Asn Cys Tyr Val Tyr Asn Lys Pro Gly Glu Asp Val Val Val Met Ala
                85                  90                  95

Gln Thr Leu Glu Lys Val Phe Leu Gln Lys Ile Glu Ser Met Pro
            100                 105                 110
```

```
<210>   22
<211>   109
<212>   PRT
<213>   Saccharomyces cerevisiae

<400>   22
```

```
Asn Pro Ile Pro Lys His Gln Gln Lys His Ala Leu Leu Ala Ile Lys
1               5                   10                  15

Ala Val Lys Arg Leu Lys Asp Ala Arg Pro Phe Leu Gln Pro Val Asp
                20                  25                  30

Pro Val Lys Leu Asp Ile Pro Phe Tyr Phe Asn Tyr Ile Lys Arg Pro
        35                  40                  45

Met Asp Leu Ser Thr Ile Glu Arg Lys Leu Asn Val Gly Ala Tyr Glu
    50                  55                  60

Val Pro Glu Gln Ile Thr Glu Asp Phe Asn Leu Met Val Asn Asn Ser
65                  70                  75                  80

Ile Lys Phe Asn Gly Pro Asn Ala Gly Ile Ser Gln Met Ala Arg Asn
                85                  90                  95

Ile Gln Ala Ser Phe Glu Lys His Met Leu Asn Met Pro
            100                 105
```

<210>   23
<211>   113
<212>   PRT
<213>   Homo sapiens

<400>   23

Lys Lys Gly Lys Leu Ser Glu Gln Leu Lys His Cys Asn Gly Ile Leu
1   5   10   15

Lys Glu Leu Leu Ser Lys Lys His Ala Ala Tyr Ala Trp Pro Phe Tyr
                20   25   30

Lys Pro Val Asp Ala Ser Ala Leu Gly Leu His Asp Tyr His Asp Ile
            35   40   45

Ile Lys His Pro Met Asp Leu Ser Thr Val Lys Arg Lys Met Glu Asn
        50   55   60

Arg Asp Tyr Arg Asp Ala Gln Glu Phe Ala Ala Asp Val Arg Leu Met
65   70   75   80

Phe Ser Asn Cys Tyr Lys Tyr Asn Pro Pro Asp His Asp Val Val Ala
                85   90   95

Met Ala Arg Lys Leu Gln Asp Val Phe Glu Phe Arg Tyr Ala Lys Met
            100   105   110

Pro

<210>   24
<211>   113
<212>   PRT
<213>   Homo sapiens

<400>   24

Lys Lys Gly Lys Leu Ser Glu His Leu Arg Tyr Cys Asp Ser Ile Leu
1   5   10   15

Arg Glu Met Leu Ser Lys Lys His Ala Ala Tyr Ala Trp Pro Phe Tyr
                20   25   30

Lys Pro Val Asp Ala Glu Ala Leu Glu Leu His Asp Tyr His Asp Ile
            35   40   45

Ile Lys His Pro Met Asp Leu Ser Thr Val Lys Arg Lys Met Asp Gly
        50   55   60

Arg Glu Tyr Pro Asp Ala Gln Gly Phe Ala Ala Asp Val Arg Leu Met
65   70   75   80

Phe Ser Asn Cys Tyr Lys Tyr Asn Pro Pro Asp His Glu Val Val Ala
                85   90   95

Met Ala Arg Lys Leu Gln Asp Val Phe Glu Met Arg Phe Ala Lys Met
            100   105   110

Pro

<210>   25
<211>   113
<212>   PRT
<213>   Drosophila melanogaster

<400>   25

```
Asn Lys Glu Lys Leu Ser Asp Ala Leu Lys Ser Cys Asn Glu Ile Leu
1               5                   10                  15

Lys Glu Leu Phe Ser Lys Lys His Ser Gly Tyr Ala Trp Pro Phe Tyr
            20                  25                  30

Lys Pro Val Asp Ala Glu Met Leu Gly Leu His Asp Tyr His Asp Ile
        35                  40                  45

Ile Lys Lys Pro Met Asp Leu Gly Thr Val Lys Arg Lys Met Asp Asn
    50                  55                  60

Arg Glu Tyr Lys Ser Ala Pro Glu Phe Ala Ala Asp Val Arg Leu Ile
65                  70                  75                  80

Phe Thr Asn Cys Tyr Lys Tyr Asn Pro Pro Asp His Asp Val Val Ala
                85                  90                  95

Met Gly Arg Lys Leu Gln Asp Val Phe Glu Met Arg Tyr Ala Asn Ile
            100                 105                 110

Pro


<210>   26
<211>   113
<212>   PRT
<213>   Saccharomyces cerevisiae

<400>   26

Lys Ser Lys Arg Leu Gln Gln Ala Met Lys Phe Cys Gln Ser Val Leu
1               5                   10                  15

Lys Glu Leu Met Ala Lys Lys His Ala Ser Tyr Asn Tyr Pro Phe Leu
            20                  25                  30

Glu Pro Val Asp Pro Val Ser Met Asn Leu Pro Thr Tyr Phe Asp Tyr
        35                  40                  45

Val Lys Glu Pro Met Asp Leu Gly Thr Ile Ala Lys Lys Leu Asn Asp
    50                  55                  60

Trp Gln Tyr Gln Thr Met Glu Asp Phe Glu Arg Glu Val Arg Leu Val
65                  70                  75                  80

Phe Lys Asn Cys Tyr Thr Phe Asn Pro Asp Gly Thr Ile Val Asn Met
                85                  90                  95

Met Gly His Arg Leu Glu Glu Val Phe Asn Ser Lys Trp Ala Asp Arg
            100                 105                 110

Pro


<210>   27
<211>   108
<212>   PRT
<213>   Homo sapiens

<400>   27

Met Glu Met Gln Leu Thr Pro Phe Leu Ile Leu Leu Arg Lys Thr Leu
1               5                   10                  15

Glu Gln Leu Gln Glu Lys Asp Thr Gly Asn Ile Phe Ser Glu Pro Val
            20                  25                  30
```

```
Pro Leu Ser Glu Val Pro Asp Tyr Leu Asp His Ile Lys Lys Pro Met
        35              40              45

Asp Phe Phe Thr Met Lys Gln Asn Leu Glu Ala Tyr Arg Tyr Leu Asn
    50              55              60

Phe Asp Asp Phe Glu Glu Asp Phe Asn Leu Ile Val Ser Asn Cys Leu
65              70              75              80

Lys Tyr Asn Ala Lys Asp Thr Ile Phe Tyr Arg Ala Ala Val Arg Leu
            85              90              95

Arg Glu Gln Gly Gly Ala Val Val Arg Gln Ala Arg
            100             105
```

```
<210>   28
<211>   113
<212>   PRT
<213>   Homo sapiens

<400>   28
```

```
Ser Glu Asp Gln Glu Ala Ile Gln Ala Gln Lys Ile Trp Lys Lys Ala
1               5               10              15

Ile Met Leu Val Trp Arg Ala Ala Ala Asn His Arg Tyr Ala Asn Val
        20              25              30

Phe Leu Gln Pro Val Thr Asp Asp Ile Ala Pro Gly Tyr His Ser Ile
        35              40              45

Val Gln Arg Pro Met Asp Leu Ser Thr Ile Lys Lys Asn Ile Glu Asn
    50              55              60

Gly Leu Ile Arg Ser Thr Ala Glu Phe Gln Arg Asp Ile Met Leu Met
65              70              75              80

Phe Gln Asn Ala Val Met Tyr Asn Ser Ser Asp His Asp Val Tyr His
            85              90              95

Met Ala Val Glu Met Gln Arg Asp Val Leu Glu Gln Ile Gln Gln Phe
            100             105             110

Leu
```

```
<210>   29
<211>   106
<212>   PRT
<213>   Gallus gallus

<400>   29
```

```
Asn Leu Pro Thr Val Asp Pro Ile Ala Val Cys His Glu Leu Tyr Asn
1               5               10              15

Thr Ile Arg Asp Tyr Lys Asp Glu Gln Gly Arg Leu Leu Cys Glu Leu
            20              25              30

Phe Ile Arg Ala Pro Lys Arg Arg Asn Gln Pro Asp Tyr Tyr Glu Val
        35              40              45

Val Ser Gln Pro Ile Asp Leu Met Lys Ile Gln Gln Lys Leu Lys Met
    50              55              60

Glu Glu Tyr Asp Asp Val Asn Val Leu Thr Ala Asp Phe Gln Leu Leu
65              70              75              80

Phe Asn Asn Ala Lys Ala Tyr Tyr Lys Pro Asp Ser Pro Glu Tyr Lys
```

```
                    85                    90                    95

Ala Ala Cys Lys Leu Trp Glu Leu Tyr Leu
            100                   105


<210>  30
<211>  112
<212>  PRT
<213>  Gallus gallus

<400>  30

Ser Ser Pro Gly Tyr Leu Lys Glu Ile Leu Glu Gln Leu Leu Glu Ala
1               5                   10                  15

Val Ala Val Ala Thr Asn Pro Ser Gly Arg Leu Ile Ser Glu Leu Phe
            20                  25                  30

Gln Lys Leu Pro Ser Lys Val Gln Tyr Pro Asp Tyr Tyr Ala Ile Ile
        35                  40                  45

Lys Glu Pro Ile Asp Leu Lys Thr Ile Ala Gln Arg Ile Gln Asn Gly
    50                  55                  60

Thr Tyr Lys Ser Ile His Ala Met Ala Lys Asp Ile Asp Leu Leu Ala
65                  70                  75                  80

Lys Asn Ala Lys Thr Tyr Asn Glu Pro Gly Ser Gln Val Phe Lys Asp
                85                  90                  95

Ala Asn Ala Ile Lys Lys Ile Phe Asn Met Lys Lys Ala Glu Ile Glu
            100                 105                 110

<210>  31
<211>  112
<212>  PRT
<213>  Gallus gallus

<400>  31

Thr Ser Phe Met Asp Thr Ser Asn Pro Leu Tyr Gln Leu Tyr Asp Thr
1               5                   10                  15

Val Arg Ser Cys Arg Asn Asn Gln Gly Gln Leu Ile Ser Glu Pro Phe
            20                  25                  30

Phe Gln Leu Pro Ser Lys Lys Lys Tyr Pro Asp Tyr Tyr Gln Gln Ile
        35                  40                  45

Lys Thr Pro Ile Ser Leu Gln Gln Ile Arg Ala Lys Leu Lys Asn His
    50                  55                  60

Glu Tyr Glu Thr Leu Asp Gln Leu Glu Ala Asp Leu Asn Leu Met Phe
65                  70                  75                  80

Glu Asn Ala Lys Arg Tyr Asn Val Pro Asn Ser Ala Ile Tyr Lys Arg
                85                  90                  95

Val Leu Lys Met Gln Gln Val Met Gln Ala Lys Lys Lys Glu Leu Ala
            100                 105                 110

<210>  32
<211>  113
<212>  PRT
<213>  Gallus gallus

<400>  32

Ser Lys Lys Asn Met Arg Lys Gln Arg Met Lys Ile Leu Tyr Asn Ala
```

```
1                 5                      10                     15
Val Leu Glu Ala Arg Glu Ser Gly Thr Gln Arg Arg Leu Cys Asp Leu
            20                  25                  30

Phe Met Val Lys Pro Ser Lys Lys Asp Tyr Pro Asp Tyr Tyr Lys Ile
        35                  40                  45

Ile Leu Glu Pro Met Asp Leu Lys Met Ile Glu His Asn Ile Arg Asn
    50                  55                  60

Asp Lys Tyr Val Gly Glu Glu Ala Met Ile Asp Asp Met Lys Leu Met
65                  70                  75                  80

Phe Arg Asn Ala Arg His Tyr Asn Glu Glu Gly Ser Gln Val Tyr Asn
            85                  90                  95

Asp Ala His Met Leu Glu Lys Ile Leu Lys Glu Lys Arg Lys Glu Leu
            100                 105                 110

Gly
```

```
<210>    33
<211>    115
<212>    PRT
<213>    Gallus gallus

<400>    33

Lys Lys Ser Lys Tyr Met Thr Pro Met Gln Gln Lys Leu Asn Glu Val
1               5                   10                  15

Tyr Glu Ala Val Lys Asn Tyr Thr Asp Lys Arg Gly Arg Arg Leu Ser
            20                  25                  30

Ala Ile Phe Leu Arg Leu Pro Ser Arg Ser Glu Leu Pro Asp Tyr Tyr
            35                  40                  45

Ile Thr Ile Lys Lys Pro Val Asp Met Glu Lys Ile Arg Ser His Met
        50                  55                  60

Met Ala Asn Lys Tyr Gln Asp Ile Asp Ser Met Val Glu Asp Phe Val
65                  70                  75                  80

Met Met Phe Asn Asn Ala Cys Thr Tyr Asn Glu Pro Glu Ser Leu Ile
            85                  90                  95

Tyr Lys Asp Ala Leu Val Leu His Lys Val Leu Leu Glu Thr Arg Arg
            100                 105                 110

Glu Ile Glu
        115
```

```
<210>    34
<211>    112
<212>    PRT
<213>    Description of unknown organism, see Jeanmougin et al., Trends
in Biochem. Sci. 22:151-153 (1997)

<400>    34

His Asn Ala Pro Phe Asp Lys Thr Lys Phe Asp Glu Val Leu Glu Ala
1               5                   10                  15

Leu Val Gly Leu Lys Asp Asn Glu Gly Asn Pro Phe Asp Asp Ile Phe
            20                  25                  30

Glu Glu Leu Pro Ser Lys Arg Tyr Phe Pro Asp Tyr Tyr Gln Ile Ile
```

```
           35                    40                    45
Gln Lys Pro Ile Cys Tyr Lys Met Met Arg Asn Lys Ala Lys Thr Gly
    50                    55                    60

Lys Tyr Leu Ser Met Gly Asp Phe Tyr Asp Asp Ile Arg Leu Met Val
65                    70                    75                    80

Ser Asn Ala Gln Thr Tyr Asn Met Pro Gly Ser Leu Val Tyr Glu Cys
                85                    90                    95

Ser Val Leu Ile Ala Asn Thr Ala Asn Ser Leu Glu Ser Lys Asp Gly
                100                   105                   110
```

```
<210>  35
<211>  113
<212>  PRT
<213>  Description of unknown organism, see Jeanmougin et al., Trends
in Biochem. Sci. 22:151-153 (1997)

<400>  35
```

```
Gly Thr Asn Glu Ile Asp Val Pro Lys Val Ile Gln Asn Ile Leu Asp
1               5                   10                  15

Ala Leu His Glu Glu Lys Asp Glu Gln Gly Arg Phe Leu Ile Asp Ile
            20                  25                  30

Phe Ile Asp Leu Pro Ser Lys Arg Leu Tyr Pro Asp Tyr Tyr Glu Ile
          35                  40                  45

Ile Lys Ser Pro Met Thr Ile Lys Met Leu Glu Lys Arg Phe Lys Lys
      50                  55                  60

Gly Glu Tyr Thr Thr Leu Glu Ser Phe Val Lys Asp Leu Asn Gln Met
65                  70                  75                  80

Phe Ile Asn Ala Lys Thr Tyr Asn Ala Pro Gly Ser Phe Val Tyr Glu
                85                  90                  95

Asp Ala Glu Lys Leu Ser Gln Leu Ser Ser Ser Leu Ile Ser Ser Phe
                100                 105                 110

Ser
```

```
<210>  36
<211>  113
<212>  PRT
<213>  Homo sapiens

<400>  36
```

```
Gly Thr Asn Glu Ile Asp Val Pro Lys Val Ile Gln Asn Ile Leu Asp
1               5                   10                  15

Ala Leu His Glu Glu Lys Asp Glu Gln Gly Arg Phe Leu Ile Asp Ile
            20                  25                  30

Phe Ile Asp Leu Pro Ser Lys Arg Leu Tyr Pro Asp Tyr Tyr Glu Ile
          35                  40                  45

Ile Lys Ser Pro Met Thr Ile Lys Met Leu Glu Lys Arg Phe Lys Lys
      50                  55                  60

Gly Glu Tyr Thr Thr Leu Glu Ser Phe Val Lys Asp Leu Asn Gln Met
65                  70                  75                  80

Phe Ile Asn Ala Lys Thr Tyr Asn Ala Pro Gly Ser Phe Val Tyr Glu
```

                    85   90   95

Asp Ala Glu Lys Leu Ser Gln Leu Ser Ser Ser Leu Ile Ser Ser Phe
            100   105   110

Ser

<210>   37
<211>   114
<212>   PRT
<213>   Homo sapiens

<400>   37

Ser Pro Asn Pro Pro Asn Leu Thr Lys Lys Met Lys Lys Ile Val Asp
1   5   10   15

Ala Val Ile Lys Tyr Lys Asp Ser Ser Ser Gly Arg Gln Leu Ser Glu
            20   25   30

Val Phe Ile Gln Leu Pro Ser Arg Lys Glu Leu Pro Glu Tyr Tyr Glu
        35   40   45

Leu Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg
    50   55   60

Asn His Lys Tyr Arg Ser Leu Asn Asp Leu Glu Lys Asp Val Met Leu
65   70   75   80

Leu Cys Gln Asn Ala Gln Thr Phe Asn Leu Glu Gly Ser Leu Ile Tyr
                85   90   95

Glu Asp Ser Ile Val Leu Gln Ser Val Phe Thr Ser Val Arg Gln Lys
            100   105   110

Ile Glu

<210>   38
<211>   113
<212>   PRT
<213>   Gallus gallus

<400>   38

Ser Pro Asn Pro Pro Lys Leu Thr Lys Gln Met Asn Ala Ile Ile Asp
1   5   10   15

Thr Val Ile Asn Tyr Lys Asp Ser Ser Gly Arg Gln Leu Ser Glu Val
            20   25   30

Phe Ile Gln Leu Pro Ser Arg Lys Glu Leu Pro Glu Tyr Tyr Glu Leu
        35   40   45

Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg Asn
    50   55   60

His Lys Tyr Arg Ser Leu Gly Asp Leu Glu Lys Asp Val Met Leu Leu
65   70   75   80

Cys His Asn Ala Gln Thr Phe Asn Leu Glu Gly Ser Gln Ile Tyr Glu
                85   90   95

Asp Ser Ile Val Leu Gln Ser Val Phe Lys Ser Ala Arg Gln Lys Ile
            100   105   110

Ala

<210> 39
<211> 114
<212> PRT
<213> Gallus gallus

<400> 39

Ser Pro Asn Pro Pro Asn Leu Thr Lys Lys Met Lys Lys Ile Val Asp
1   5   10   15

Ala Val Ile Lys Tyr Lys Asp Ser Ser Ser Gly Arg Gln Leu Ser Glu
            20   25   30

Val Phe Ile Gln Leu Pro Ser Arg Lys Glu Leu Pro Glu Tyr Tyr Glu
        35   40   45

Leu Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg
    50   55   60

Asn His Lys Tyr Arg Ser Leu Asn Asp Leu Glu Lys Asp Val Met Leu
65   70   75   80

Leu Cys Gln Asn Ala Gln Thr Phe Asn Leu Glu Val Ser Leu Ile Tyr
                85   90   95

Glu Asp Ser Ile Val Leu Gln Ser Val Phe Thr Ser Val Arg Gln Lys
            100   105   110

Ile Glu

<210> 40
<211> 105
<212> PRT
<213> Homo sapiens

<400> 40

Ala Lys Leu Ser Pro Ala Asn Gln Arg Lys Cys Glu Arg Val Leu Leu
1   5   10   15

Ala Leu Phe Cys His Glu Pro Cys Arg Pro Leu His Gln Leu Ala Thr
            20   25   30

Asp Ser Thr Phe Ser Leu Asp Gln Pro Gly Gly Thr Leu Asp Leu Thr
            35   40   45

Leu Ile Arg Ala Arg Leu Gln Glu Lys Leu Ser Pro Pro Tyr Ser Ser
    50   55   60

Pro Gln Glu Phe Ala Gln Asp Val Gly Arg Met Phe Lys Gln Phe Asn
65   70   75   80

Lys Leu Thr Glu Asp Lys Ala Asp Val Gln Ser Ile Ile Gly Leu Gln
                85   90   95

Arg Phe Phe Glu Thr Arg Met Asn Glu
            100   105

<210> 41
<211> 105
<212> PRT
<213> Mus musculus

<400> 41

Ala Lys Leu Ser Pro Ala Asn Gln Arg Lys Cys Glu Arg Val Leu Leu
1   5   10   15

```
Ala Leu Phe Cys His Glu Pro Cys Arg Pro Leu His Gln Leu Ala Thr
            20              25              30

Asp Ser Thr Phe Ser Met Glu Gln Pro Gly Gly Thr Leu Asp Leu Thr
            35              40              45

Leu Ile Arg Ala Arg Leu Gln Glu Lys Leu Ser Pro Pro Tyr Ser Ser
    50              55              60

Pro Gln Glu Phe Ala Gln Asp Val Gly Arg Met Phe Lys Gln Phe Asn
65              70              75              80

Lys Leu Thr Glu Asp Lys Ala Asp Val Gln Ser Ile Ile Gly Leu Gln
                85              90              95

Arg Phe Phe Glu Thr Arg Met Asn Asp
            100             105
```

```
<210>   42
<211>   108
<212>   PRT
<213>   Mus sp.

<400>   42
```

```
Thr Lys Leu Thr Pro Ile Asp Lys Arg Lys Cys Glu Arg Leu Leu Leu
1               5               10              15

Phe Leu Tyr Cys His Glu Met Ser Leu Ala Phe Gln Asp Pro Val Pro
            20              25              30

Leu Thr Val Pro Asp Tyr Tyr Lys Ile Ile Lys Asn Pro Met Asp Leu
            35              40              45

Ser Thr Ile Lys Lys Arg Leu Gln Glu Asp Tyr Cys Met Tyr Thr Lys
    50              55              60

Pro Glu Asp Phe Val Ala Asp Phe Arg Leu Ile Phe Gln Asn Cys Ala
65              70              75              80

Glu Phe Asn Glu Pro Asp Ser Glu Val Ala Asn Ala Gly Ile Lys Leu
                85              90              95

Glu Ser Tyr Phe Glu Glu Leu Leu Lys Asn Leu Tyr
            100             105
```

```
<210>   43
<211>   14
<212>   PRT
<213>   artificial sequence

<220>
<221>   X
<222>   (1)..(2)
<223>   X can be any single amino acid


<220>
<221>   X
<222>   (4)..(4)
<223>   X is two to three amino acids. Each of these can be any amino
aci
        d


<220>
<221>   X
<222>   (6)..(6)
```

<223>   X is five to eight amino acids. Each of these can be any amino
ac
        id

<220>
<221>   X
<222>   (7)..(7)
<223>   X is a single amino acid that can be Pro, Lys, or His.

<220>
<221>   X
<222>   (8)..(8)
<223>   X is a single amino acid that can be any amino acid.

<220>
<221>   X
<222>   (10)..(10)
<223>   X is a single amino acid that can be a Tyr, Phe, or His.

<220>
<221>   X
<222>   (11)..(11)
<223>   X is five amino acids. Each of these can be any amino acid.

<220>
<221>   X
<222>   (13)..(13)
<223>   X is a single amino acid that can be Met, Ile, or Val.

<400>   43

Xaa Xaa Phe Xaa Pro Xaa Xaa Xaa Tyr Xaa Xaa Pro Xaa Asp
1   5   10

<210>   44
<211>   20
<212>   PRT
<213>   artificial sequence

<400>   44

Trp Pro Phe Met Glu Pro Val Lys Arg Thr Glu Ala Pro Gly Tyr Tyr
1   5   10   15

Glu Val Ile Arg
            20

<210>   45
<211>   101
<212>   PRT
<213>   Human immunodeficiency virus type 1

<400>   45

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser
1   5   10   15

Gln Pro Lys Thr Ala Ser Asn Asn Cys Tyr Cys Lys Arg Cys Cys Leu
            20   25   30

His Cys Gln Val Cys Phe Thr Lys Lys Gly Leu Gly Ile Ser Tyr Gly
        35   40   45

```
Arg Lys Lys Arg Arg Gln Arg Arg Arg Ala Pro Gln Asp Ser Lys Thr
    50              55              60
His Gln Val Ser Leu Ser Lys Gln Pro Ala Ser Gln Pro Arg Gly Asp
65              70              75              80
Pro Thr Gly Pro Lys Glu Ser Lys Lys Lys Val Glu Arg Glu Thr Glu
                85              90              95
Thr Asp Pro Glu Asp
            100
```

```
<210>  46
<211>  7
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is one to three amino acids.  Each amino acid can be any amino
       acid


<400>  46

Tyr Gly Arg Lys Xaa Arg Gln
1               5

<210>  47
<211>  10
<212>  PRT
<213>  artificial sequence

<400>  47

Ser Tyr Gly Arg Lys Lys Arg Arg Gln Arg
1               5               10

<210>  48
<211>  10
<212>  PRT
<213>  artificial

<220>
<221>  X
<222>  (2)..(2)
<223>  X is two to four amino acids.  Each of these can be any amino
aci
       d


<220>
<221>  X
<222>  (4)..(4)
<223>  X is two to four amino acids.  Each of these can be any amino
aci
       d


<220>
<221>  X
<222>  (6)..(6)
<223>  X is two to four amino acids.  Each of these can be any amino
aci
       d


<220>
```

```
<221>  X
<222>  (8)..(8)
<223>  X is one to three amino acids.  Each of these can be any amino
ac
       id


<220>
<221>  X
<222>  (10)..(10)
<223>  X is a single amino acid that is either Ile, Leu, Met, or Val.


<400>  48

Phe Xaa Val Xaa Glu Xaa Tyr Xaa Val Xaa
1               5                   10


<210>  49
<211>  62
<212>  PRT
<213>  artificial sequence

<400>  49

Phe Met Glu Pro Val Lys Arg Thr Glu Ala Pro Gly Tyr Tyr Glu Val
1               5                   10                  15

Ile Arg Phe Pro Met Asp Leu Lys Thr Met Ser Glu Arg Leu Lys Asn
            20                  25                  30

Arg Tyr Tyr Val Ser Lys Lys Leu Phe Met Ala Asp Leu Gln Arg Val
        35                  40                  45

Phe Thr Asn Cys Lys Glu Tyr Asn Ala Ala Glu Ser Glu Tyr
    50                  55                  60


<210>  50
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK).


<400>  50

Ser Tyr Gly Arg Xaa Lys Arg Arg Gln Arg Cys
1               5                   10


<210>  51
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK)


<400>  51

Ser Ala Gly Arg Xaa Lys Arg Arg Gln Arg Cys
1               5                   10
```

```
<210>  52
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X ia an acetylated lysine (AcK)


<400>  52

Ser Tyr Gly Ala Xaa Lys Arg Arg Gln Arg Cys
1               5                   10

<210>  53
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK).


<400>  53

Ser Tyr Gly Arg Xaa Ala Arg Arg Gln Arg Cys
1               5                   10

<210>  54
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK).


<400>  54

Ser Tyr Gly Arg Xaa Lys Ala Arg Gln Arg Cys
1               5                   10

<210>  55
<211>  11
<212>  PRT
<213>  artificial sequence

<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK)


<400>  55

Ser Tyr Gly Arg Xaa Lys Arg Ala Gln Arg Cys
1               5                   10

<210>  56
<211>  11
<212>  PRT
<213>  artificial sequence
```

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<220>
<221>  X
<222>  (5)..(5)
<223>  X is an acetylated lysine (AcK)


<400>  56

Ser Tyr Gly Arg Xaa Lys Arg Arg Ala Arg Cys
1                 5                 10

<210>  57
<211>  11
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<220>
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<400>  57

Ser Tyr Gly Arg Lys Xaa Arg Arg Gln Arg Cys
1                 5                 10

<210>  58
<211>  11
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<220>
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<400>  58

Thr Asn Cys Tyr Cys Lys Xaa Cys Cys Phe His
1                 5                 10

<210>  59
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<220>
<221>  X
<222>  (16)..(16)
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<400>  59

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1                 5                 10                15

Arg His Arg Lys
            20
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